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CLAIMS

- 1. A metabolically engineered micro-organism having an operative first metabolic pathway in which a first 5 metabolite is transformed into a second metabolite in a reaction in which NAD is a cofactor for a first enzyme, said reaction step producing NADH, and in which said second metabolite is transformed into at least one further metabolite in a reaction catalysed by a second 10 enzyme, and having an operative second metabolic pathway characterised by an enzyme activity in excess of a native level in respect of a third enzyme catalysing a non-reversible reaction in which NADP is a cofactor and NADPH is a product and in which said first 15 metabolite is transformed into a said further metabolite without the involvement of said second enzyme.
- 2. A micro-organism as claimed in claim 1, wherein said first metabolic pathway is a native pathway.
 - 3. A micro-organism as claimed in any preceding claim, wherein said first enzyme is a phosphorylating dehydrogenase.

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- 4. A micro-organism as claimed in claim 1 or claim 2, wherein said second enzyme is a kinase.
- 5. A micro-organism as claimed in claim 3, wherein said third enzyme is a non-phosphorylating dehydrogenase.

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- 6. A micro-organism as claimed in claim 5, wherein said third enzyme is GAPN (EC 1.2.1.9).
- 7. A micro-organism as claimed in claim 6, wherein said first enzyme is GAPDH (EC 1.2.1.12).
 - 8. A micro-organism as claimed in any preceding claim, wherein at least one copy of a genetic sequence encoding said third enzyme has been recombinantly introduced into said organism.
- A micro-organism as claimed in any preceding claim,
 wherein a genetic sequence encoding said third enzyme
 is operatively linked to an expression signal not
 natively associated with said genetic sequence in said
 organism.
 - 10. A micro-organism as claimed in any preceding claim which is a yeast.

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11. A micro-organism as claimed in claim 10, which is a micro-organism belonging to the genus Saccharomyces, Klyuveromyces, Candida, Pichia, Debaromyces, Hansenula, Yarrowia, Zygosaccharomyces or Schizosaccharomyces.

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12. A micro-organism as claimed in claim 10, which is a strain of Saccharomyces cerevisiae, S. kluyveri, S. bayanus, S. exiguus, S. sevazzi, S. uvarum, Klyuveromyces lactis K. marxianus var. marxianus, K. thermotolerans, Candida utilis C. tropicalis, Pichia stipidis, P. pastoris, P. sorbitophila, Debaromyces hansenii, Hansenula polymorpha, Yarrowia lipolytica,

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Zygosaccharomyces rouxii or Schizosaccharomyces pombe..

13. A genetically transformed micro-organism containing one or more copies of an heterologous DNA sequence encoding GAPN operatively associated with an expression signal and having a functional native or heterologous expression capability for GAPDH (EC 1.2.12).

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- 14. A method of producing a desired metabolic product with decreased production of an undesired metabolic product, comprising culturing a micro-organism as claimed in any preceding claim.
- 15. A method as claimed in claim 14, wherein the desired
 15 product is ethanol, lactic acid, citric acid, an amino acid or an antibiotic.
- 16. A method as claimed in claim 14 or claim 15, wherein said undesired metabolic product is glycerol, acetate or an amino acid.